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OM protein - protein search, using sw model

Run on: February 1, 2005, 14:23:28 ; Search time 147 Seconds
(without alignments)
594.776 Million cell updates/sec

Title: US-10-629-329A-2
Perfect score: 1322
Sequence: 1 MSGCDAGEGDCCRRCAQD.....SMKKVGLDPSQLPVGNGIV 242

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1608061 seqs, 361289386 residues

Total number of hits satisfying chosen parameters: 1608061

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	ID	Description
1	1322	100.0	242	US-10-629-329A-2	Sequence 2, Appli
2	1307	98.9	242	US-10-220-381-2	Sequence 2, Appli
3	1296	98.0	242	US-10-381-710-4	Sequence 4, Appli
4	1239.5	93.8	241	US-10-381-710-2	Sequence 2, Appli
5	1239.5	93.8	241	US-10-629-329A-4	Sequence 4, Appli
6	573.5	43.4	529	US-10-437-963-195546	Sequence 195546
7	563	42.6	256	US-10-424-599-157170	Sequence 157170
8	563	42.6	517	US-10-425-115-253963	Sequence 253963
9	563	42.6	524	US-10-425-114-64486	Sequence 64486, A
10	555	42.0	497	US-10-425-114-65135	Sequence 65135, A
11	555	41.8	517	US-10-767-701-45914	Sequence 45914, A
12	549	41.5	522	US-10-424-599-273717	Sequence 273717
13	549	41.5	540	US-10-425-114-46271	Sequence 46271, A

14	514.5	38.9	594	17	US-10-425-115-253964	Sequence 253964,
15	467.5	35.4	459	15	US-10-425-114-61505	Sequence 61505, A
16	415	31.4	533	15	US-10-425-114-57875	Sequence 57875, A
17	312	23.6	64	14	US-10-029-386-28983	Sequence 28983, A
18	292	22.1	128	16	US-10-767-701-59226	Sequence 59226, A
19	228	17.2	59	14	US-10-106-698-5148	Sequence 5148, Ap
20	214	16.2	202	14	US-10-156-761-14187	Sequence 14187, A
21	195	14.8	212	15	US-10-282-122A-45375	Sequence 45375, A
22	193	14.6	204	15	US-10-282-122A-69836	Sequence 69836, A
23	182	13.8	204	15	US-10-282-122A-55860	Sequence 55860, A
24	166	12.6	205	15	US-10-282-122A-43563	Sequence 43563, A
25	163.5	12.4	241	14	US-10-156-761-13948	Sequence 13948, A
26	145	11.0	231	10	US-09-557-796-32	Sequence 32, Appl
27	141	10.7	104	15	US-10-424-599-183312	Sequence 183312,
28	140	10.6	202	15	US-10-282-122A-59882	Sequence 59882, A
29	137	10.4	234	15	US-10-282-122A-74190	Sequence 74190, A
30	137	10.4	238	10	US-09-557-796-18	Sequence 18, Appl
31	136.5	10.3	220	15	US-10-282-122A-49784	Sequence 49784, A
32	133.5	10.1	234	15	US-10-282-122A-74354	Sequence 74354, A
33	132	10.0	228	15	US-10-282-122A-75285	Sequence 75285, A
34	131	9.9	181	15	US-10-282-122A-46808	Sequence 46808, A
35	131	9.9	228	15	US-10-282-122A-76130	Sequence 76130, A
36	129.5	9.8	231	10	US-09-557-796-33	Sequence 33, Appl
37	129	9.8	228	15	US-10-282-122A-56798	Sequence 56798, A
38	128.5	9.7	231	15	US-10-282-122A-77900	Sequence 77900, A
39	127.5	9.6	232	15	US-10-282-122A-57636	Sequence 57636, A
40	127	9.6	230	15	US-10-282-122A-77598	Sequence 77598, A
41	125.5	9.5	236	15	US-10-282-122A-57141	Sequence 57141, A
42	122.5	9.3	228	15	US-10-282-122A-53018	Sequence 53018, A
43	118	8.9	213	15	US-10-282-122A-50759	Sequence 50759, A
44	114.5	8.7	212	15	US-10-282-122A-48161	Sequence 48161, A
45	112.5	8.5	233	15	US-10-282-122A-51793	Sequence 51793, A

ALIGNMENTS

RESULT 1

US-10-629-329A-2
; Sequence 2, Application US/10629329A
; Publication No. US2004008648A1
; GENERAL INFORMATION:
; APPLICANT: DARNAY, BRYANT G.
; TITLE OF INVENTION: METHODS AND COMPOSITIONS USING POLYNUCLEOTIDES AND
; FILE OF INVENTION: POLYPEPTIDES OF RANK-ASSOCIATED INHIBITOR (RAIN)
; FILE REFERENCE: UTSC:761US
; CURRENT APPLICATION NUMBER: US/10/629,329A
; CURRENT FILING DATE: 2003-07-29
; PRIOR APPLICATION NUMBER: 60/399,205
; PRIOR FILING DATE: 2002-07-29
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 242
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-629-329A-2

Query Match	100.0%	Score 1322;	DB 15;	Length 242;
Best Local Similarity	100.0%	Pred. No. 4,6e-129;		
Matches 242;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	MSGCDAGEGDCCRRCAQDKHEPRYLIPELCKQPYHGLGWVTGTGGISLKHGDEIYIAP	60	
Db	1	MSGCDAGEGDCCRRCAQDKHEPRYLIPELCKQPYHGLGWVTGTGGISLKHGDEIYIAP	60	
Qy	61	SGVQKERTQPEDMFVCDINEKDISGPSKSLKKSQCTPLFNAYTMRGAGAVIHTHSKA	120	
Db	61	SGVQKERTQPEDMFVCDINEKDISGPSKSLKKSQCTPLFNAYTMRGAGAVIHTHSKA	120	
Qy	121	AVMATLLPFGREFKITHQEMIKGIKCTSGGYRVDMLVPIENTPEKGLKDRMAHA	180	
Db	121	AVMATLLPFGREFKITHQEMIKGIKCTSGGYRVDMLVPIENTPEKGLKDRMAHA	180	

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QY 181 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db 181 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
QY 241 IV 242
Db 241 IV 242

RESULT 2
US-10-220-381-2
; Sequence 2, Application US/10220381
; Publication No. US20030207430A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.
; APPLICANT: TANG, Y. Tom
; APPLICANT: LU, Dying Aina M.
; APPLICANT: BANDMAN, Olga
; APPLICANT: YUE, Henry
; APPLICANT: AZIMZAI, Yalda
; APPLICANT: LAL, Preeti
; APPLICANT: BURFORD, Neil
; APPLICANT: BAUGHN, Mariah R.
; TITLE OF INVENTION: HUMAN ENZYME MOLECULES
; FILE REFERENCE: PF-0763 PCT
; CURRENT APPLICATION NUMBER: US/10/220,381
; CURRENT FILING DATE: 2001-03-01
; NUMBER OF SEQ ID NOS: 52
; SOFTWARE: PERL Program
; SEQ ID NO 2
; LENGTH: 242
; TYPE: PRT
; ORGANISM: Homo sapiens
; NAME/KEY: misc feature
; FEATURE:
; OTHER INFORMATION: Incyte ID No. US20030207430A1 2116390CD1
; US-10-220-381-2

Query Match 98.9%; Score 1307; DB 14; Length 242;
Best Local Similarity 99.2%; Pred. No. 1.7e-127;
Matches 240; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db 1 MSGCDAREGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
QY 121 AVMATLLFPGRFVKITHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db 121 AVMATLLFPGRFVKITHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
QY 181 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db 181 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
QY 241 IV 242
Db 241 IV 242

RESULT 3
US-10-381-710-4
; Sequence 4, Application US/10381710
; Publication No. US20040052789A1
; GENERAL INFORMATION:
; APPLICANT: SHA, Shiken et al.
; TITLE OF INVENTION: NOVEL PROTEINS, GENES ENCODING THEM AND METHOD OF USING THE SAME
; FILE REFERENCE: 0230-0198P
; CURRENT APPLICATION NUMBER: US/10/381,710
; CURRENT FILING DATE: 2003-09-16
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: Patent in version 3.2
; SEQ ID NO 2
; LENGTH: 241
; TYPE: PRT
; ORGANISM: Mouse macrophage cell RAW 264.7
; US-10-381-710-2

Query Match 93.8%; Score 1239.5; DB 15; Length 241;
Best Local Similarity 93.8%; Pred. No. 1.8e-120;
Matches 227; Conservative 9; Mismatches 5; Indels 1; Gaps 1;

QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db 1 MSGCQA-QGDCCSRFCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 59
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db 60 SGVQKERIQPEDMFVCDINEQDISGPPASKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 119
QY 121 AVMATLLFPGRFVKITHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db 120 AVMATLLFPGRFVKITHQEMIKIRKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 179
QY 181 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db 180 MNEYPDSCAVLVRHGVVWGETWEKAKTMCECYDYLFDIAVSMKKVGLDPTQLPVGENG 239
QY 241 IV 242
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Db 240 IV 241

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RESULT 5

US-10-629-329A-4

; Sequence 4, Application US/10629329A

; Publication No. US20040086848A1

; GENERAL INFORMATION:

; APPLICANT: DARWAY, BRYANT G.

; TITLE OF INVENTION: METHODS AND COMPOSITIONS USING POLYNUCLEOTIDES AND

; TITLE OF INVENTION: POLYPEPTIDES OF RANK-ASSOCIATED INHIBITOR (RAIN)

; FILE REFERENCE: UTS:761US

; CURRENT APPLICATION NUMBER: US/10/629,329A

; BEST FILING DATE: 2003-07-29

; PRIOR APPLICATION NUMBER: 60/399,205

; FILING DATE: 2002-07-29

; NUMBER OF SEQ ID NOS: 23

; SOFTWARE: Patent In Ver. 2.1

; SEQ ID NO 4

; LENGTH: 241

; TYPE: PRT

; ORGANISM: Mus musculus

US-10-629-329A-4

Query Match 93.8%; Score 1239.5; DB 15; Length 241;

Best Local Similarity 93.8%; Pred. No. 1.8e-120;

Matches 227; Conservative 9; Mismatches 5; Indels 1; Gaps 1;

Qy 1 MSGDAGEGDCSRGCAQDKHPRYLIPELCKQFYHLGWVTGTTGGGSLKHGDEIYIAP 60

Db 1 MSGCQA-QGDCCSRCAQDKHPRFLIPELCKQFYHLGWVTGTTGGGSLKHGNEIYIAP 59

Qy 61 SGVOKERIQPDMFVCDINEKDISGPPSKLKSCTPLEMNAITMRGAGAVIHTSKA 120

Db 60 SGVOKERIQPDMFVCDINEKDISGPPSKLKSCTPLEMNAITMRGAGAVIHTSKA 119

Qy 121 AVMATLLPFGREFKITHQEMIKGIKCTSGGYRYDDMLVPIENTPEEKGLKDRMAHA 180

Db 120 AVMATLLPFGREFKITHQEMIKGIKCTSGGYRYDDMLVPIENTPEEKGLKDRMAHA 179

Qy 181 MNEYPDSCAVLVRHGVYWGWTWEKATMCBCYDLFDIAVSMKKVGLDPSQLPVGENG 240

Db 180 MNEYPDSCAVLVRHGVYWGWTWEKATMCBCYDLFDIAVSMKKOMGLDPTQLPVGENG 239

Qy 241 IV 242

Db 240 IV 241

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RESULT 6

US-10-437-963-195546

; Sequence 195546, Application US/10437963

; Publication No. US20040123343A1

; GENERAL INFORMATION:

; APPLICANT: La Rosa, Thomas J.

; APPLICANT: Kovalic, David K.

; APPLICANT: Zhou, Yihua

; APPLICANT: Cao, Yongwei

; APPLICANT: Wu, Wei

; APPLICANT: Boukharov, Andrey A.

; APPLICANT: Barbazuk, Brad

; APPLICANT: Li, Ping

; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With

; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

; FILE REFERENCE: 38-21(53221)B

; CURRENT APPLICATION NUMBER: US/10/437,963

; FILING DATE: 2003-05-14

; NUMBER OF SEQ ID NOS: 204966

; SEQ ID NO 195546

; LENGTH: 529

; TYPE: PRT

; ORGANISM: Oryza sativa

us-10-629-329a-2.rapb

Wed Feb 2 10:19:19 2005

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Best Local Similarity 47.0%; Pred. No. 1.7e-49;
Matches 118; Conservative 31; Mismatches 68; Indels 34; Gaps 6;

QY 2 SGCDAGEGDCSRRCGAODKE-----HPRYLIPELCKQFYHLGWVTGTGGGISLKH 52
DB 11 SGCS-----CEAAVGAMASEAYLEGAPVREARELVAELCRHFYAGQWVTGTGGSITVKV 64
QY 53 GDE-----IYIAPSGVQKRIQPEDMFVCDINEKDISGPS-----PSKLLKKSQCTPL 100
DB 65 NDPVPLADRLIIVMSPSGVQKRMVADGMVMAADGKVLAPVAKPWPKNPKCTDCAPL 124
QY 101 FNNAYTMRGAGAVIHTHSHKAAVMATLLFPG-REFKITHQEMIKGIKCKTSGGYRYDDML 159
DB 125 FMKAYLMRGAGAVIHTSHGICTIATMLIPGAKFRVTHMEMIKGIK-----HGYHDEL 178
QY 160 VVPIIENPTPEEKGLKDRMAHAMNEYDPDSCAVLVRHGGVYVWGETWEKAKTMCBCYDLYFD 219
DB 179 VIPIIENPTPEYELTDSLSIAIAYPKATAVLVRNHGIYVWGESWINAKTQAECHYLLD 238
QY 220 IAVSMKKVGLD 230
DB 239 ACIKLYQLGID 249

RESULT 10
US-10-425-114-65135
; Sequence 65135, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E.
; APPLICANT: Tabaska, Jack E.
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 65135
; LENGTH: 497
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-ZMFLMO17103D03_FLI.pep
US-10-425-114-65135
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Query Match 42.0%; Score 555; DB 15; Length 497;
Best Local Similarity 50.2%; Pred. No. 1.1e-48;
Matches 114; Conservative 32; Mismatches 61; Indels 20; Gaps 5;

QY 17 GAODKEHPRYLIPELCKQFYHLGWVTGTGGGISLKHGDE-----IYIAPSGVQKRI 68
DB 3 GAPVRE-ARELVAELCRHFYAGQWVTGTGGSITVKVNDPAVPLADRLIIVMSPSGVQKRM 61
QY 69 QPEDMFVCDINEKDISGPS-----PSKLLKKSQCTPLFNNAYTMRGAGAVIHTHSHKAAVMA 124
DB 62 VVADMTVMAADGKVLAPVAKPWPKNPKCTDCAPLFMKAYLMRGAGAVIHTSHGICTCIA 121
QY 125 TLLFPG-REFKITHQEMIKGIKCKTSGGYRYDDMLVVPPIIENPTPEEKGLKDRMAHAMNE 183
DB 122 TMLIPGAKFRVTHMEMIKGIK-----HGYHDELVPIIENPTPEYELTDSLSIAIAA 175
QY 184 YPDSCAVLVRHGGVYVWGETWEKAKTMCBCYDLYFDIAVSMKKVGLD 230
DB 176 YPKATAVLVRNHGIYVWGESWINAKTQAECHYLLDACCILKLYQLGID 222

RESULT 11
US-10-767-701-45914
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US-10-425-115-253963
; Sequence 253963, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 253963
; LENGTH: 517
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_163196C.1.pep
US-10-425-115-253963
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Query Match 42.6%; Score 563; DB 17; Length 517;
Best Local Similarity 47.0%; Pred. No. 1.7e-49;
Matches 118; Conservative 31; Mismatches 68; Indels 34; Gaps 6;

QY 2 SGCDAGEGDCSRRCGAODKE-----HPRYLIPELCKQFYHLGWVTGTGGGISLKH 52
DB 4 SGCS-----CEAAVGAMASEAYLEGAPVREARELVAELCRHFYAGQWVTGTGGSITVKV 57
QY 53 GDE-----IYIAPSGVQKRIQPEDMFVCDINEKDISGPS-----PSKLLKKSQCTPL 100
DB 58 NDPVPLADRLIIVMSPSGVQKRMVADGMVMAADGKVLAPVAKPWPKNPKCTDCAPL 117
QY 101 FNNAYTMRGAGAVIHTHSHKAAVMATLLFPG-REFKITHQEMIKGIKCKTSGGYRYDDML 159
DB 118 FMKAYLMRGAGAVIHTSHGICTIATMLIPGAKFRVTHMEMIKGIK-----HGYHDEL 171
QY 160 VVPIIENPTPEEKGLKDRMAHAMNEYDPDSCAVLVRHGGVYVWGETWEKAKTMCBCYDLYFD 219
DB 172 VIPIIENPTPEYELTDSLSIAIAYPKATAVLVRNHGIYVWGESWINAKTQAECHYLLD 231
QY 220 IAVSMKKVGLD 230
DB 232 ACIKLYQLGID 242
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RESULT 9
US-10-425-114-64486
; Sequence 64486, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E.
; APPLICANT: Tabaska, Jack E.
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 64486
; LENGTH: 524
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB3060-104-F8_FLI.pep
US-10-425-114-64486
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; Sequence 45914, Application US/10767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(533535)B
; CURRENT APPLICATION NUMBER: US/10/767,701
; CURRENT FILING DATE: 2004-01-29
; NUMBER OF SEQ ID NOS: 63128
; SEQ ID NO 45914
; LENGTH: 517
; TYPE: PRT
; ORGANISM: Sorghum bicolor
; FEATURE:
; OTHER INFORMATION: Clone ID: SORBI-28MAY03-C2601_1.pep
US-10-767-701-45914

Query Match 41.8%; Score 553; DB 16; Length 517;
Best Local Similarity 46.1%; Pred. No. 1.8e-48;
Matches 117; Conservative 34; Mismatches 71; Indels 32; Gaps 7;

Qy 9 GDC-CSRRCQAQDK-----HPRYLIPELCKQFYHLGWVTGGTGGISLKHGDE--- 55
Db 4 GCGCEAAVAGTASEAYLEGEVPREARELVAELCRHFAQGWVTGGTGGITVKVNDPVP 63

Qy 56 ----IYIAPSGVQKRIQPEDMFVCDINEKDISGPS-----PSKKLKKSQCTPLFMNAYT 106
Db 64 LADRLVWSPSGVQKRIQPEDMFVCDINEKDISGPS-----PSKKLKKSQCTPLFMNAYT 123

Qy 107 MRGAGAVIHTHSKAAVMATLLPFG-REFKITHQEMIKGKCTGGYRYDDMLVVPPIE 165
Db 124 MRGAGAVIHTSHGEMETCIATMLNPGAKERFWTHMEMIKGK-----HGYRDELVIPIVE 177

Qy 166 NTPPEKGLKDRMAHMANEYPSDCAVLVRHGVYVNGETWEKATWCBCYDYLFDIAVSMK 225
Db 178 NTPPEYELTSLSEIAAAYPKATAVLVRNHGIVYVNGDSWINAKTQAECHYLLDACIKLY 237

Qy 226 KVGLED---PSQLPV 236
Db 238 QLGIDWTTPEHGPI 251

RESULT 12
US-10-424-599-273717
; Sequence 273717, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J
; APPLICANT: Kovalic, David K
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 273717
; LENGTH: 522
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_89188C.1.pep
US-10-424-599-273717

Query Match 41.5%; Score 549; DB 15; Length 522;
Best Local Similarity 51.3%; Pred. No. 4.9e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGTGGISLKHGDE-----IYIAPSGVQKRIQPEDMFVC 76

RESULT 14
US-10-425-115-253964
; Sequence 253964, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 46271
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701207771_FLI.pep
US-10-425-114-46271

Query Match 41.5%; Score 549; DB 15; Length 540;
Best Local Similarity 51.3%; Pred. No. 5.1e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGTGGISLKHGDE-----IYIAPSGVQKRIQPEDMFVC 76
Db 47 RALMAELCRHFYTLGWVTGGTGGISLKHGDESDIPRQQLILMAPSGVQKRIQPEDMFVC 106

Qy 77 DINEKDISGPS-----KKLKKSQCTPLFMNAYTMRGAGAVIHTHSKAAVMATLLP-GR 131
Db 107 SHSGSVLSAPSPKWPHPKPCDCDPLFKKAYENRDAAVFHSHGIESCLVTMINPLSK 166

Qy 132 EFKITHQEMIKGKCTGGYRYDDMLVPIENTPEKGLKDRMAHMANEYPSDCAVL 191
Db 167 EFRITHMEMIKGK---GHGY---DELVVPPIENTAYEYQLTESFAKAIEDYPKATAVL 220

Qy 192 VRRHGVYVNGETWEKATWCBCYDYLFDIAVSMKKVGLD---PSQLPV 236
Db 221 VRNHGVFVNGDSWISAKTQSECYHYLFDAALKHQMGLEDWSTPNHGPI 268

RESULT 18
US-10-425-114-46271
; Sequence 46271, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 46271
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701207771_FLI.pep
US-10-425-114-46271

Query Match 41.5%; Score 549; DB 15; Length 540;
Best Local Similarity 51.3%; Pred. No. 5.1e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGTGGISLKHGDE-----IYIAPSGVQKRIQPEDMFVC 76
Db 47 RALMAELCRHFYTLGWVTGGTGGISLKHGDESDIPRQQLILMAPSGVQKRIQPEDMFVC 106

Qy 77 DINEKDISGPS-----KKLKKSQCTPLFMNAYTMRGAGAVIHTHSKAAVMATLLP-GR 131
Db 107 SHSGSVLSAPSPKWPHPKPCDCDPLFKKAYENRDAAVFHSHGIESCLVTMINPLSK 166

Qy 132 EFKITHQEMIKGKCTGGYRYDDMLVPIENTPEKGLKDRMAHMANEYPSDCAVL 191
Db 167 EFRITHMEMIKGK---GHGY---DELVVPPIENTAYEYQLTESFAKAIEDYPKATAVL 220

Qy 192 VRRHGVYVNGETWEKATWCBCYDYLFDIAVSMKKVGLD---PSQLPV 236
Db 221 VRNHGVFVNGDSWISAKTQSECYHYLFDAALKHQMGLEDWSTPNHGPI 268

RESULT 18
US-10-425-115-253964
; Sequence 253964, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 46271
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701207771_FLI.pep
US-10-425-114-46271

Query Match 41.5%; Score 549; DB 15; Length 540;
Best Local Similarity 51.3%; Pred. No. 5.1e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGTGGISLKHGDE-----IYIAPSGVQKRIQPEDMFVC 76
Db 47 RALMAELCRHFYTLGWVTGGTGGISLKHGDESDIPRQQLILMAPSGVQKRIQPEDMFVC 106

Qy 77 DINEKDISGPS-----KKLKKSQCTPLFMNAYTMRGAGAVIHTHSKAAVMATLLP-GR 131
Db 107 SHSGSVLSAPSPKWPHPKPCDCDPLFKKAYENRDAAVFHSHGIESCLVTMINPLSK 166

Qy 132 EFKITHQEMIKGKCTGGYRYDDMLVPIENTPEKGLKDRMAHMANEYPSDCAVL 191
Db 167 EFRITHMEMIKGK---GHGY---DELVVPPIENTAYEYQLTESFAKAIEDYPKATAVL 220

Qy 192 VRRHGVYVNGETWEKATWCBCYDYLFDIAVSMKKVGLD---PSQLPV 236
Db 221 VRNHGVFVNGDSWISAKTQSECYHYLFDAALKHQMGLEDWSTPNHGPI 268
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Wed Feb 2 10:19:19 2005

112 AVIHTHKAAMATLPPG-REFKITHOEMIKGIKCTSGGYRYDDMLVVPPIENTPEE 170
71 AVIHTSHGIECTIATMLIPGAKEPRVTHMEMIKIGK-----HGYHDELVIPIENTPVE 124
171 KGLKDRMAHAMNEYPDSCAVLVRHGHVYVWGGETWEKAKTMCBCYDYLFDIAVSMKKVGLD 230
125 YELTDSLSEALAAYPKATAVLVRNHGIYVWGESWINAKTQACFYHLLDACIKLYQLGID 184

Search completed: February 1, 2005, 14:33:32
Job time : 149 secs

FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425.115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 253964
LENGTH: 594
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_163197C.1.pap
US-10-425-115-253964

Query Match 38.9%; Score 514.5; DB 17; Length 594;
Best Local Similarity 36.0%; Pred. No. 2.3e-44;
Matches 118; Conservative 31; Mismatches 68; Indels 111; Gaps 7;

QY 2 SGCDAGEGDCSRRRCQAQDKE-----HPRVLIPELCKQFYHLGWVTGTGGGISLKH 52
DB 4 SGCS-----CEAAVGAASEAYLEGAPVREARELVAELCRHFAQGWVTGTGGSIYKV 57
QY 53 GDE-----IYIAPSGVOKERIOPEDMFVCDINEKDISGPS-----PSKKLKSQCTPL 100
DB 58 NDPTVPLADRLIVMSPSGVOKERWVAEDMYMAADGKVLAPVAKPWPKPKCTDCAPL 117
QY 101 FMNAYTMRGAGAVIHTHKAAMATLPPG-REFKITHOEMIKGIKCTSGGYRYDDML 159
DB 118 FMKAYLMRGAGAVIHTSHGIECTIATMLIPGAKEFRVTHMEMIKIGK-----HGYHDEL 171
QY 160 VPIIENTPBEKGLKDRMAHAMNEYPDSCAVLVRHGHVYVWGGETWEKAKT----- 210
DB 172 VPIIENTPYEYELTDSLSEALAAYPKATAVLVRNHGIYVWGESWINAKTQACFGFRDQ 231
QY 211 ----- 210
DB 232 IKDFIWTLPKEPDLHASFRKNTMYLHIFMGSLILQSFVIIITDVLVAGGIWGRNS 291
QY 211 -----CECYDYLFDIAMSKVGLD 230
DB 292 LTPAFCREACFYHLLDACIKLYQLGID 319

RESULT 15
US-10-425-114-61505
Sequence 61505, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425.114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 61505
LENGTH: 459
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
OTHER INFORMATION: Clone ID: LIB143-005-F3_FLI.pap
US-10-425-114-61505

Query Match 35.4%; Score 467.5; DB 15; Length 459;
Best Local Similarity 51.7%; Pred. No. 1.3e-39;
Matches 93; Conservative 27; Mismatches 49; Indels 11; Gaps 3;
QY 56 IYIAPSGVOKERIOPEDMFVCDINEKDISGPS-----PSKKLKSQCTPLFMNAYTMRGAG 111
DB 11 IVMSPSGVOKERWVAEDMYMAADGKVLAPVAKPWPKPKCTDCAPLFMKAYLMRGAG 70